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OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/003,035

DATE: 12/11/2001

TIME: 11:47:43

Input Set : A:\Genphar.712.txt

Output Set: N:\CRF3\12112001\I003035.raw

ENTERED

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3 <110> APPLICANT: Wang, Danher
5 <120> TITLE OF INVENTION: GENETIC VACCINE AGAINST HUMAN IMMUNODEFICIENCY VIRUS
7 <130> FILE REFERENCE: 22488-712
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/003,035
C--> 9 <141> CURRENT FILING DATE: 2001-11-01
9 <150> PRIOR APPLICATION NUMBER: 09/585,599
10 <151> PRIOR FILING DATE: 2000-06-02
12 <150> PRIOR APPLICATION NUMBER: PCT/US01/18238
13 <151> PRIOR FILING DATE: 2001-06-04
15 <160> NUMBER OF SEQ ID NOS: 75
17 <170> SOFTWARE: PatentIn version 3.1
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20 <211> LENGTH: 7
21 <212> TYPE: DNA
22 <213> ORGANISM: Ebola virus
24 <400> SEQUENCE: 1
25 tttttttt 7
28 <210> SEQ ID NO: 2
29 <211> LENGTH: 7
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31 <213> ORGANISM: Ebola virus
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34 uuuuuuuu 7
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38 <211> LENGTH: 8
39 <212> TYPE: RNA
40 <213> ORGANISM: Artificial sequence
42 <220> FEATURE:
43 <223> OTHER INFORMATION: Modified RNA editing site
45 <400> SEQUENCE: 3
46 uucuucuu 8
49 <210> SEQ ID NO: 4
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51 <212> TYPE: DNA
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54 <400> SEQUENCE: 4
55 aaaaaaaa 7
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60 <212> TYPE: DNA
61 <213> ORGANISM: Artificial sequence
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64 <223> OTHER INFORMATION: DNA of modified RNA editing site
66 <400> SEQUENCE: 5
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70 <210> SEQ ID NO: 6
71 <211> LENGTH: 6

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73 <213> ORGANISM: Ebola virus
75 <400> SEQUENCE: 6
76 uuuuuu 6
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82 <213> ORGANISM: Artificial sequence
84 <220> FEATURE:
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87 <400> SEQUENCE: 7
88 uucuuu 6
91 <210> SEQ ID NO: 8
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93 <212> TYPE: DNA
94 <213> ORGANISM: Artificial sequence
96 <220> FEATURE:
97 <223> OTHER INFORMATION: DNA of modified RNA editing site
99 <400> SEQUENCE: 8
100 ttcttc 6
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104 <211> LENGTH: 21
105 <212> TYPE: PRT
106 <213> ORGANISM: Homo sapiens
108 <400> SEQUENCE: 9
110 Ser Trp Leu Leu Leu Leu Leu Ser Leu Ser Leu Leu Gln Ala Thr
111 1 5 10 15
114 Asp Phe Met Ser Leu
115 20
118 <210> SEQ ID NO: 10
119 <211> LENGTH: 5
120 <212> TYPE: PRT
121 <213> ORGANISM: Ebola virus
123 <400> SEQUENCE: 10
125 Arg Arg Thr Arg Arg
126 1 5
129 <210> SEQ ID NO: 11
130 <211> LENGTH: 4
131 <212> TYPE: PRT
132 <213> ORGANISM: Human immunodeficiency virus type 1
134 <400> SEQUENCE: 11
136 Arg Glu Lys Arg
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140 <210> SEQ ID NO: 12
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142 <212> TYPE: DNA
143 <213> ORGANISM: Artificial sequence
145 <220> FEATURE:
146 <223> OTHER INFORMATION: DNA of GAG site

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148 <400> SEQUENCE: 12
149 ggagctggt
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154 <212> TYPE: PRT
155 <213> ORGANISM: Artificial sequence
157 <220> FEATURE:
158 <223> OTHER INFORMATION: GAG site
160 <400> SEQUENCE: 13
162 Gly Ala Gly
163 1
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169 <213> ORGANISM: Artificial sequence
171 <220> FEATURE:
172 <223> OTHER INFORMATION: Env/Tat/Rev
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179 gaagcatcca ggaagtcagc ctaaaactgc ttgtaccaat tgctattgta aaaagtgttg 180
181 ctttcattgc caagtttggt tcataacaaa agccttaggc atctcctatg gcaggaagaa 240
183 ggcggagacag cgacgaagac ctctcaagg cagtcagact catcaagttt ctctatcaaa 300
185 gcagtaagta gtacatgtaa tgcaacctat acaaatagca atagtagcat tagtagtagc 360
187 aataataata gcaatagtgt tgtggtccat agtaatcata gaatatagga aaatattaag 420
189 acaaagaaaa atagacaggt taattgatag actaatagaa agagcagaag acagtggcaa 480
191 tgagagtga ggaagaaat cagcacttgt ggagatgggg gtggagatgg ggcaccatgc 540
193 tccttgggat gttgatgatc tgtagtgtca cagaaaaatt gtgggtcaca gtctattatg 600
195 ggggtacctgt gtggaaggaa gcaaccacca ctctattttg tgcatacagat gctaaagcat 660
197 atgatacaga ggtacataat gtttgggcca cacatgcctg tgtaccaca gacccaacc 720
199 cacaagaagt agtattggt aatgtgacag aaaattttta catgtggaaa aatgacatgg 780
201 tagaacagat gcatgaggat ataatcagtt tatgggatca aagcctaaag ccatgtgtaa 840
203 aattaacccc actctgtgtt agtttaagt gcaatgattt gaagaatgat actaatacca 900
205 atagtagtag cgggagaatg ataattggaga aaggagagat aaaaaactgc tctttcaata 960
207 tcagcacaag cataagaggt aaggtgcaga aagaatatgc atttttttat aaacttgata 1020
209 taataccaat agataatgat actaccagct atacgttgac aagttgtaac acctcagtca 1080
211 ttacacaggc ctgtccaaag gtatcctttg agccaattcc catacattat tgtgccccgg 1140
213 ctggttttgc gattctaaaa tgtaataata agacgttcaa tggaacagga ccatgtacaa 1200
215 atgtcagcac agtacaatgt acacatggaa ttaggccagt agtatcaact caactgctgt 1260
217 taaatggcag tctggcgaaa gaagaggtag taattagatc tgccaatttc acagacaatg 1320
219 ctaaaacccat aatagtacag ctgaaccaat ctgtagaaat taattgtaca agaccaaca 1380
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223 gaaaaatagg aaatatgaga caagcacatt gtaacattag tagagcaaaa tgggaataaca 1500
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227 ttaagcagtc ctcaggaggg gaccagaaa ttgtaacgca cagttttaat tgtggagggg 1620
229 aatttttcta ctgtaattca acacaactgt ttaatagtag ttggtttaat agtacttgga 1680
231 gtactaaagg gtcaaataac actgaaggaa gtgacacaat caccctccca tgcagaataa 1740
233 aacaaattat aaacatgtgg caggaagtag gaaaagcaat gtatgccctt cccatcagtg 1800
235 gacaaattag atgttcacat aatattacag ggctgctatt aacaagagat ggtggttaata 1860

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237 gcaacaatga gtccgagatc ttcagacctg gaggaggaga tatgagggac aattggagaa 1920
239 gtgaattata taaatataaa gtagtaaaaa ttgaaccatt aggagtagca cccaccaagg 1980
241 caaagagaag agtgggtgcag agagaaaaaa gagcagtggg aataggagct ttgttccttg 2040
243 ggttccttggg agcagcagga agcactatgg gcgcagcgtc aatgacgctg acgggtacagg 2100
245 ccagacaatt attgtctggt atagtgcagc agcagaacaa tttgctgagg gctattgagg 2160
247 cgcaacagca tctgttgcaa ctccacagtct ggggcatcaa gcagctccag gcaagaatcc 2220
249 tggctgtgga aagataccta aaggatcaac agctcctggg gatttggggg tgctctggaa 2280
251 aactcatttg caccactgct gtgccttgga atgctagtgg gagtaataaa tctctggaac 2340
253 agatttggaa taacatgacc tggatggagt gggacagaga aattaacaat tacacaagct 2400
255 taatacactc cttaattgaa gaatcgcaaa accagcaaga aaagaatgaa caagaattat 2460
257 tgggaattaga taaatgggca agtttgtgga attgggttta cataacaaat tggctgtggg 2520
259 atataaaatt attcataatg atagtaggag gcttggtagg ttaagaata gtttttgctg 2580
261 tactttctgt agtgaataga gttaggcagg gatattcacc attatcgttt cagacccacc 2640
263 tcccaatccc gaggggaccc gacaggcccg aaggaataga agaagaaggg ggagagagag 2700
265 acagagacag atccattcga ttagtgaacg gatcccttagc acttatctgg gacgatctgc 2760
267 ggagcctgtg cctcttcagc taccaccgct tgagagactt actcttgatt gtaacgagga 2820
269 ttgtggaact tctgggacgc agggggtggg aagccctcaa atattggtgg aatctcctac 2880
271 agtattggag tcaggagcta aagaatagtg ctgttagctt gctcaatgcc acagctatag 2940
273 cagtagctga ggggacagat aggggttatag aagtagtaca aggagcttat agagctattc 3000
275 gccacatacc tagaagaata agacagggct tggaaaggat tttgctataa gatgggtggc 3060
277 aagtgggtcaa aaagtgtgtg ggttggatgg cctgctgtaa gggaaagaat gagacgagct 3120
279 gagccagcag cagatggggg gggagcagca tctcgag 3157

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282 <210> SEQ ID NO: 15

283 <211> LENGTH: 508

284 <212> TYPE: DNA

285 <213> ORGANISM: Artificial sequence

287 <220> FEATURE:

288 <223> OTHER INFORMATION: Modified IL-2

290 <400> SEQUENCE: 15

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291 tcaactctct taactactac tcacagtaac ctcaactcct gccacaatgt acaggatgca 60
293 actcctgtct tgcattgcac taagtcttgc acttgtcaca aacagtgcac ctacttcaag 120
295 ttctacaaaag aaaacacagc tacaactgga gcatttactg ctggatttac agatgatttt 180
297 gaatggaatt aataattaca agaatcccaa actcaccagg atgctcacat ttaagtttta 240
299 catgcccaag aaggccacag aactgaaaca tcttcagtgt cttgaagaag aactcaaacc 300
301 tctggaggaa gtgctaaatt tagctcaaag caaaaacttt cacttaagac ccagggactt 360
303 aatcagcaat atcaacgtaa tagttctgga actaaaggga tctgaaacaa cattcatgtg 420
305 tgaatatgct gatgagacag caaccattgt agaatttctg aacagatgga ttaccttttg 480
307 tcaaagcatc atctcaacac taacttga 508

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310 <210> SEQ ID NO: 16

311 <211> LENGTH: 2280

312 <212> TYPE: DNA

313 <213> ORGANISM: Artificial sequence

315 <220> FEATURE:

316 <223> OTHER INFORMATION: Modified Env

318 <400> SEQUENCE: 16

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319 gaattcgcca ccatgggagt gaaggagaaa tatcagcact tgtggagatg ggggtggaga 60
321 tggggcacca tgctccttgg gatgttgatg atctgtagtg ctacagaaaa attgtgggtc 120
323 acagtctatt atgggggtacc tgtgtggaag gaagcaacca ccaactctatt ttgtgcatca 180
325 gatgctaaag catatgatac agaggtagat aatgttttgg ccacacatgc ctgtgtaccc 240

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329 aaaaatgaca tggtagaaca gatgcatgag gatataatca gtttatggga tcaaagccta 360
331 aagccatgtg taaaattaac cccactctgt gttagtttta agtgactga tttgaagaat 420
333 gatactaata ccaatagtag tagcgggaga atgataatgg agaaaggaga gataaaaaac 480
335 tgctctttca atatcagcac aagcataaga ggtaagggtg agaaagaata tgcatttttt 540
337 tataaacttg atataatacc aatagataat gatactacca gctatacgtt gacaagttgt 600
339 aacacctcag tcattacaca ggctgtcca aaggtatcct ttgagccaat tcccatacat 660
341 tattgtgccc cggctggttt tgcgattcta aaatgtaata ataagacgtt caatggaaca 720
343 ggacctgtga caaatgtcag cacagtacaa tgtacacatg gaattaggcc agtagtatca 780
345 actcaactgc tgtaaagtgg cagtctggca gaagaagagg tagtaattag atctgccaat 840
347 ttcacagaca atgctaaaac cataatagta cagctgaacc aatctgtaga aattaattgt 900
349 acaagaccca acaacaatac aagaaaaagt atccgtatcc agagaggacc agggagagca 960
351 tttgttacia taggaaaaat aggaaatatg agacaagcac attgtaacat tagtagagca 1020
353 aaatggaata acactttaaa acagatagat agcaaattaa gagaacaatt tggaaataat 1080
355 aaaacaataa tctttaagca gtctcagga ggggaccag aaattgtaac gcacagtttt 1140
357 aattgtggag gggaattttt ctactgtaat tcaacacaac tgtttaatag tacttggttt 1200
359 aatagtactt ggagtactaa aggttcaa atactgaag gaagtgcac aatcaccctc 1260
361 ccatgcagaa taaaacaaat tataaacatg tggcaggaag taggaaaagc aatgtatgcc 1320
363 cctcccatca gtggacaaat tagatgttca tcaaataatta cagggtctgt attaacaaga 1380
365 gatggtggtg atagcaacaa tgagtccgag atcttcagac ctggaggagg agatatgagg 1440
367 gacaattgga gaagtgaatt atataaatat aaagtagtaa aaattgaacc attaggagta 1500
369 gcacccacca aggcaaagag aagagtgggt cagactagtg cagtgggaat aggagctttg 1560
371 ttccttggtt tcttgaggc agcaggaagc actatggcg cagcgtcaat gacgtgacg 1620
373 gtacaggcca gacaattatt gtctggtata gtgcagcagc agaacaattt gctgagggtc 1680
375 attgaggcgc aacagcatct gttgcaactc acagtctggg gcatcaagca gctccaggca 1740
377 agaatcctgg ctgtggaaag atacctaaag gatcaacagc tcctggggat ttgggggttg 1800
379 tctggaaaac tcatttgcac cactgctgtg ccttggaatg ctagtggag taataaatct 1860
381 ctggaacaga tttggaataa catgacctgg atggagtggg acagagaaat taacaattac 1920
383 acaagcttaa tacactcctt aattgaagaa tcgcaaaacc agcaagaaaa gaatgaacaa 1980
385 gaattatttg aattagataa atgggcaagt ttgtggaatt ggtttaacat aacaaattgg 2040
387 ctgtggtata taaaattatt cataatgata gtaggaggct tggtaggttt aagaatagtt 2100
389 tttgctgtac tttctgtagt gaatagagtt aggcagggat attcaccatt atcgtttcag 2160
391 acccacctcc caatcccag gggacccgac aggcccgag gaatagaaga agaagggtga 2220
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397 <211> LENGTH: 1496
398 <212> TYPE: DNA
399 <213> ORGANISM: Artificial sequence
401 <220> FEATURE:
402 <223> OTHER INFORMATION: Full length Gag
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407 gatgggaaaa aattcggtta aggccagggg gaaagaaaaa atataaatta aaacatatag 120
409 tatgggcaag caggagcta gaacgactac aaccatccct tcagacagga tcagaagaac 180
411 ttagatcatt atataatata gtagcaaccc tctattgtgt gcatcaaagg atagagataa 240
413 aagacaccaa ggaagcttta gacaagatag aggaagagca aaacaaaagt aagaaaaaag 300
415 cacagcaagc agcagctgac acaggacaca gcagtcaggc cagccaaaat taccctatag 360
417 tgcagaacat ccaggggcaa atggtacatc aggccatata acctagaact ttaaatgcat 420
419 gggtaaaagt agtagaagag aaggctttca gccagaagt aatacccatg ttttcagcat 480

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VERIFICATION SUMMARY

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L:9 M:270 C: Current Application Number differs, Replaced Current Application No

L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date